OIPE 1205

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RAW SEQUENCE LISTING PATENT APPLICATION US/09/865,989

First 5 pages)

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1			SEQUENCE LISTING
2			
3	(1)	GENERAL	INFORMATION:
4			
5		(i)	APPLICANT: Dasseux, Jean-Louis
6			Sekul, Renate
7			Buttner, Klaus
8			Cornut, Isabelle
9			Metz, Gunther
10			
11		(ii)	TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
12		(11)	AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
13			AND INDIK OUD TO INDIK DIDDITIDDING DIDORDING
14		(444)	NUMBER OF SEQUENCES: 258
15		(111)	NOMBER OF BEQUENCES. 256
		()	CODDEGDONDENGE ADDREGG
16			CORRESPONDENCE ADDRESS:
17			ADDRESSEE: Pennie & Edmonds LLP
18		, ,	STREET: 1155 Avenue of the Americas
19		, ,	CITY: New York STATE: NY COUNTRY: USA ENTERED
20			STATE: NY
21			
22		(F)	ZIP: 10036-2811
23			
24		(v) C	COMPUTER READABLE FORM:
25			MEDIUM TYPE: Diskette
26		(B)	COMPUTER: IBM Compatible
27		(C)	OPERATING SYSTEM: DOS
28		(D)	SOFTWARE: FastSEQ Version 2.0
29			
30		(vi)	CURRENT APPLICATION DATA:
31		(A)	APPLICATION NUMBER:US/09/865,989
32			FILING DATE: 25-MAY-2001
33		(C)	CLASSIFICATION:
34			
35		(vii)	PRIOR APPLICATION DATA:
36			APPLICATION NUMBER: 09/465,719
37		1 1	FILING DATE:
38		(-,	
39			
40			
41		(szi i i	.) ATTORNEY/AGENT INFORMATION:
42			NAME: Coruzzi, Laura A
43			REGISTRATION NUMBER: 30,742
44		(C)	REFERENCE/DOCKET NUMBER: 009196-0006-999
45		()	TELECOMMINICATION INFORMATION.
46			TELECOMMUNICATION INFORMATION:
47			TELEPHONE: 650-493-4935
48			TELEFAX: 650-493-5556
49		(C)	TELEX: 66141 PENNIE
50			

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51	
52	(2) INFORMATION FOR SEQ ID NO:1:
53	
54	(i) SEQUENCE CHARACTERISTICS:
55	(A) LENGTH: 22 amino acids
56	(B) TYPE: amino acid
57	(C) STRANDEDNESS: single
58	(D) TOPOLOGY: linear
59	
60	(ii) MOLECULE TYPE: None
61	(ix) FEATURE:
62	,,
63	(A) NAME/KEY: Other
64	(B) LOCATION: 16
65	(D) OTHER INFORMATION: Xaa = Naphthylalanine
66	(2) Olizik Intolkaritok. Mad - Naphenylatanine
67	
68	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
69	(AI) BEGERREE BEBERREETION. BEG ID NO.1.
70	Pro Val Leu Asp Leu Phe Arg Glu Leu Leu Asn Glu Leu Leu Glu Xaa
71	1 5 10 15
72	Leu Lys Gln Lys Leu Lys
73	20
74	
75	(2) INFORMATION FOR SEQ ID NO:2:
76	(a) International Policy In No. 11.
77	(i) SEQUENCE CHARACTERISTICS:
78	(A) LENGTH: 23 amino acids
79	(B) TYPE: amino acid
80	(C) STRANDEDNESS: single
81	(D) TOPOLOGY: linear
82	(2) 10102001. 1111001
83	(ii) MOLECULE TYPE: None
84	(==, ::=================================
85	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
86	(, 1.2 1.1
87	Gly Val Leu Asp Leu Phe Arg Glu Leu Leu Asn Glu Leu Leu Glu Ala
88	1 5 10 15
89	Leu Lys Gln Lys Leu Lys Lys
90	20
91	
92	(2) INFORMATION FOR SEQ ID NO:3:
93	
94	(i) SEQUENCE CHARACTERISTICS:
95	(A) LENGTH: 22 amino acids
96	(B) TYPE: amino acid
97	(C) STRANDEDNESS: single
98	(D) TOPOLOGY: linear
99	(=, ===================================
00	(ii) MOLECULE TYPE: None

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```
101
 102
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
 103
 104
       Pro Val Leu Asp Leu Phe Arg Glu Leu Leu Asn Glu Leu Leu Glu Trp
 105
       1
                       5
 106
       Leu Lys Gln Lys Leu Lys
 107
                   20
 108
 109
                (2) INFORMATION FOR SEQ ID NO:4:
110
111
             (i) SEQUENCE CHARACTERISTICS:
112
               (A) LENGTH: 22 amino acids
113
               (B) TYPE: amino acid
114
               (C) STRANDEDNESS: single
115
               (D) TOPOLOGY: linear
116
117
            (ii) MOLECULE TYPE: None
118
119
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
120
121
       Pro Val Leu Asp Leu Phe Arg Glu Leu Leu Asn Glu Leu Leu Glu Ala
122
       1
                                           10
123
       Leu Lys Gln Lys Leu Lys
124
                  20
125
126
                (2) INFORMATION FOR SEQ ID NO:5:
127
128
            (i) SEQUENCE CHARACTERISTICS:
129
               (A) LENGTH: 23 amino acids
130
               (B) TYPE: amino acid
131
               (C) STRANDEDNESS: single
132
              (D) TOPOLOGY: linear
133
134
            (ii) MOLECULE TYPE: None
135
            (ix) FEATURE:
136
137
                (A) NAME/KEY: Other
138
                (B) LOCATION: 1
139
                (D) OTHER INFORMATION: Xaa = D-Pro
140
141
142
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
143
144
      Xaa Val Leu Asp Leu Phe Arg Glu Leu Leu Asn Glu Leu Leu Glu Ala
145
                                           10
146
      Leu Lys Gln Lys Leu Lys Lys
147
148
149
               (2) INFORMATION FOR SEQ ID NO:6:
150
```

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151	(i) SEQUENCE CHARACTERISTICS:
152	(A) LENGTH: 22 amino acids
153	(B) TYPE: amino acid
154	(C) STRANDEDNESS: single
155	(D) TOPOLOGY: linear
156	(b) Topologi. Illieat
157	(ii) MOLEGALE TURE W
	(ii) MOLECULE TYPE: None
158	(ix) FEATURE:
159	
160	(A) NAME/KEY: Other
161	(B) LOCATION: 13
162	(D) OTHER INFORMATION: Xaa = Aib
163	
164	
165	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
166	22 15 No. 0.
167	Pro Val Leu Asp Leu Phe Arg Glu Leu Leu Asn Glu Xaa Leu Glu Ala
168	1
169	Leu Lys Gln Lys Leu Lys
170	20
171	20
172	(2) INFORMATION FOR GROUP NO. 5
173	(2) INFORMATION FOR SEQ ID NO:7:
174	(i) decoupling out to describe a
175	(i) SEQUENCE CHARACTERISTICS:
	(A) LENGTH: 22 amino acids
176	(B) TYPE: amino acid
177	(C) STRANDEDNESS: single
178	(D) TOPOLOGY: linear
179	
180	(ii) MOLECULE TYPE: None
181	
182	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
183	
184	Pro Val Leu Asp Leu Phe Lys Glu Leu Leu Asn Glu Leu Leu Glu Ala
185	1 5 10 15
186	Leu Lys Gln Lys Leu Lys
187	20
188	
189	(2) INFORMATION FOR SEQ ID NO:8:
190	
191	(i) SEQUENCE CHARACTERISTICS:
192	(A) LENGTH: 22 amino acids
193	(B) TYPE: amino acid
194	(C) STRANDEDNESS: single
195	(D) TOPOLOGY: linear
196	(D) IOPOLOGI: IIHear
196	(ii) MOLECULE TURE Non-
197	(ii) MOLECULE TYPE: None
	(wi) GEOVENOR DEGERATOR
199	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
200	

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201 202	Pro Val Leu Asp Leu Phe Arg Glu Leu Leu Asn Glu Gly Leu Glu Ala 1 5 10 15
203 204	Leu Lys Gln Lys Leu Lys
205	
206	(2) INFORMATION FOR SEQ ID NO:9:
207	(1) Intolation for Big ID No.5.
208	(i) SEQUENCE CHARACTERISTICS:
209	(A) LENGTH: 22 amino acids
210	(B) TYPE: amino acid
211	(C) STRANDEDNESS: single
212	(D) TOPOLOGY: linear
213	(=) ===================================
214	(ii) MOLECULE TYPE: None
215	(==, ==================================
216	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
217	,
218	Pro Val Leu Asp Leu Phe Arg Glu Leu Gly Asn Glu Leu Leu Glu Ala
219	1 5 10 15
220	Leu Lys Gln Lys Leu Lys
221	20
222	
223	(2) INFORMATION FOR SEQ ID NO:10:
224	~
225	(i) SEQUENCE CHARACTERISTICS:
226	(A) LENGTH: 22 amino acids
227	(B) TYPE: amino acid
228	(C) STRANDEDNESS: single
229	(D) TOPOLOGY: linear
230	
231	(ii) MOLECULE TYPE: None
232	(ix) FEATURE:
233	
234	(A) NAME/KEY: Other
235	(B) LOCATION: 17
236	(D) OTHER INFORMATION: Xaa = Naphthylalanine
237	
238	
239	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:
240	
241	Pro Val Leu Asp Leu Phe Arg Glu Leu Leu Asn Glu Leu Leu Glu Ala
242	1 5 10 15
243	Xaa Lys Gln Lys
244	20
245	(0)
2 4 6 2 4 7	(2) INFORMATION FOR SEQ ID NO:11:
24 / 248	(i) GEOUPHGE GUADAGERTAGE
248 249	(i) SEQUENCE CHARACTERISTICS:
250	(A) LENGTH: 22 amino acids
	(B) TYPE: amino acid